



OIPE

RAW SEQUENCE LISTING

DATE: 01/27/2003

PATENT APPLICATION: US/10/005,480A

TIME: 10:07:02

Input Set : D:\51158-20062.txt

Output Set: N:\CRF4\01242003\J005480A.raw

4 <110> APPLICANT: Agensys
 5 Challita-Eid, Pia M.
 6 Raitano, Arthur B.
 7 Faris, Mary
 8 Hubert, Rene S.
 9 Morrison, Karen Jane Meyrick
 10 Jakobovits, Aya
 12 <120> TITLE OF INVENTION: Nucleic Acid and Corresponding Protein
 13 Entitled 161P2F10B Useful in Treatment and Detection of
 14 Cancer
 16 <130> FILE REFERENCE: 51158-20062.00
 18 <140> CURRENT APPLICATION NUMBER: US 10/005,480A
 19 <141> CURRENT FILING DATE: 2001-11-07
 21 <160> NUMBER OF SEQ ID NOS: 765
 23 <170> SOFTWARE: FastSEQ for Windows Version 4.0

Does Not Comply
Corrected Diskette Needed

raw

ERRORED SEQUENCES

8421 <210> SEQ ID NO: 765
 8422 <211> LENGTH: 871
 8423 <212> TYPE: PRT
 8424 <213> ORGANISM: Homo Sapien
 8426 <400> SEQUENCE: 765
 8427 Met Glu Ser Thr Leu Thr Leu Ala Thr Glu Gln Pro Val Lys Lys Asn
 8428 1 5 10 15
 8429 Thr Leu Lys Lys Tyr Lys Ile Ala Cys Ile Val Leu Leu Ala Leu Leu
 8430 20 25 30
 8431 Val Ile Met Ser Leu Gly Leu Gly Leu Gly Leu Gly Leu Arg Lys Leu
 8432 35 40 45
 8433 Glu Lys Gln Gly Ser Cys Arg Lys Lys Cys Phe Asp Ala Ser Phe Arg
 8434 50 55 60
 8435 Gly Leu Glu Asn Cys Arg Cys Asp Val Ala Cys Lys Asp Arg Gly Asp
 8436 65 70 75 80
 8437 Cys Cys Trp Asp Phe Glu Asp Thr Cys Val Glu Ser Thr Arg Ile Trp
 8438 85 90 95
 8439 Met Cys Asn Lys Phe Arg Cys Gly Glu Thr Arg Leu Glu Ala Ser Leu
 8440 100 105 110
 8441 Cys Ser Cys Ser Asp Asp Cys Leu Gln Lys Lys Asp Cys Cys Ala Asp
 8442 115 120 125
 8443 Tyr Lys Ser Val Cys Gln Gly Glu Thr Ser Trp Leu Glu Glu Asn Cys
 8444 130 135 140
 8445 Asp Thr Ala Gln Gln Ser Gln Cys Pro Glu Gly Phe Asp Leu Pro Pro

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RAW SEQUENCE LISTING

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Input Set : D:\51158-20062.txt

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8446	145				150				155				160
8447	Val	Ile	Leu	Phe	Ser	Met	Asp	Gly	Phe	Arg	Ala	Glu	Tyr
8448					165				170				175
8449	Trp	Asp	Thr	Leu	Met	Pro	Asn	Ile	Asn	Lys	Leu	Lys	Thr
8450				180					185				190
8451	His	Ser	Lys	Tyr	Met	Arg	Ala	Met	Tyr	Pro	Thr	Lys	Thr
8452			195					200				205	
8453	His	Tyr	Thr	Ile	Val	Thr	Gly	Leu	Tyr	Pro	Glu	Ser	His
8454		210					215				220		
8455	Asn	Asn	Met	Tyr	Asp	Val	Leu	Asn	Lys	Asn	Phe	Ser	Leu
8456	225				230					235			240
8457	Glu	Gln	Asn	Pro	Ala	Trp	Trp	His	Gly	Gln	Pro	Met	Trp
8458				245					250				255
8459	Met	Tyr	Gln	Gly	Leu	Lys	Ala	Ala	Thr	Tyr	Phe	Trp	Pro
8460			260					265				270	
8461	Val	Ala	Ile	Asn	Gly	Ser	Phe	Pro	Ser	Ile	Tyr	Met	Pro
8462			275					280				285	
8463	Ser	Val	Pro	Phe	Glu	Glu	Arg	Ile	Ser	Thr	Leu	Leu	Lys
8464		290					295				300		
8465	Leu	Pro	Lys	Ala	Glu	Arg	Pro	Arg	Phe	Tyr	Thr	Met	Tyr
8466	305				310					315			320
8467	Pro	Asp	Ser	Ser	Gly	His	Ala	Gly	Gly	Pro	Val	Ser	Ala
8468				325					330				335
8469	Lys	Ala	Leu	Gln	Val	Val	Asp	His	Ala	Phe	Gly	Met	Leu
8470			340					345				350	
8471	Leu	Lys	Gln	Arg	Asn	Ile	Leu	His	Asn	Arg	Val	Asn	Ile
8472		355					360				365		
8473	Ala	Asp	His	Gly	Met	Asp	Gln	Thr	Tyr	Cys	Asn	Lys	Met
8474		370				375				380			
8475	Thr	Asp	Tyr	Phe	Pro	Arg	Ile	Asn	Phe	Phe	Tyr	Met	Tyr
8476	385				390					395			400
8477	Ala	Pro	Arg	Ile	Arg	Ala	His	Asn	Ile	Pro	His	Asp	Phe
8478				405				410				415	
8479	Asn	Ser	Glu	Glu	Ile	Val	Arg	Asn	Leu	Ser	Cys	Arg	Lys
8480			420					425				430	
8481	His	Phe	Lys	Pro	Tyr	Leu	Thr	Pro	Asp	Leu	Pro	Lys	Arg
8482			435					440				445	
8483	Ala	Lys	Asn	Val	Arg	Ile	Asp	Lys	Val	His	Leu	Phe	Val
8484		450				455				460			
8485	Trp	Leu	Ala	Val	Arg	Ser	Lys	Ser	Asn	Thr	Asn	Cys	Gly
8486	465				470					475			480
8487	His	Gly	Tyr	Asn	Asn	Glu	Phe	Arg	Ser	Met	Glu	Ala	Ile
8488				485				490					495
8489	His	Gly	Pro	Ser	Phe	Lys	Glu	Lys	Thr	Glu	Val	Glu	Pro
8490			500					505				510	
8491	Ile	Glu	Val	Tyr	Asn	Leu	Met	Cys	Asp	Leu	Leu	Arg	Ile
8492			515					520				525	
8493	Pro	Asn	Asn	Gly	Thr	His	Gly	Ser	Leu	Asn	His	Leu	Leu
8494		530					535					540	

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```

8495 Phe Tyr Glu Pro Ser His Ala Glu Glu Val Ser Lys Phe Ser Val Cys
8496 545                               550                               555                               560
8497 Gly Phe Ala Asn Pro Leu Pro Thr Glu Ser Leu Asp Cys Phe Cys Pro
8498                               565                               570                               575
8499 His Leu Gln Asn Ser Thr Gln Leu Glu Gln Val Asn Gln Met Leu Asn
8500                               580                               585                               590
8501 Leu Thr Gln Glu Glu Ile Thr Ala Thr Val Lys Val Asn Leu Pro Phe
8502                               595                               600                               605
8503 Gly Arg Pro Arg Val Leu Gln Lys Asn Val Asp His Cys Leu Leu Tyr
8504                               610                               615                               620
8505 His Arg Glu Tyr Val Ser Gly Phe Gly Lys Ala Met Arg Met Pro Met
8506 625                               630                               635                               640
8507 Trp Ser Ser Tyr Thr Val Pro Gln Leu Gly Asp Thr Ser Pro Leu Pro
8508                               645                               650                               655
8509 Pro Thr Val Pro Asp Cys Leu Arg Ala Asp Val Arg Val Pro Pro Ser
8510                               660                               665                               670
8511 Glu Ser Gln Lys Cys Ser Phe Tyr Leu Ala Asp Lys Asn Ile Thr His
8512                               675                               680                               685
8513 Gly Phe Leu Tyr Pro Pro Ala Ser Asn Arg Thr Ser Asp Ser Gln Tyr
8514                               690                               695                               700
8515 Asp Ala Leu Ile Thr Ser Asn Leu Val Pro Met Tyr Glu Glu Phe Arg
8516 705                               710                               715                               720
8517 Lys Met Trp Asp Tyr Phe His Ser Val Leu Leu Ile Lys His Ala Thr
8518                               725                               730                               735
8519 Glu Arg Asn Gly Val Asn Val Val Ser Gly Pro Phe Asp Tyr Asn Tyr
8520                               740                               745                               750
8521 Asp Gly His Phe Asp Ala Pro Asp Glu Ile Thr Lys His Leu Ala Asn
8522                               755                               760                               765
8523 Thr Asp Val Pro Ile Pro Thr His Tyr Phe Val Val Leu Thr Ser Cys
8524                               770                               775                               780
8525 Lys Asn Ser His Thr Pro Glu Asn Cys Pro Gly Trp Leu Asp Val Leu
8526 785                               790                               795                               800
8527 Pro Phe Ile Ile Pro His Arg Pro Thr Asn Val Glu Ser Cys Pro Glu
8528                               805                               810                               815
8529 Gly Lys Pro Glu Ala Leu Trp Val Glu Glu Arg Phe Thr Ala His Ile
8530                               820                               825                               830
8531 Ala Arg Val Arg Asp Val Glu Leu Leu Thr Gly Leu Asp Phe Tyr Gln
8532                               835                               840                               845
8533 Asp Lys Val Gln Pro Val Ser Glu Ile Leu Gln Leu Lys Thr Tyr Leu
8534                               850                               855                               860
8535 Pro Thr Phe Glu Thr Thr Ile
8536 865                               870

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E--> 8540 (1)
E--> 8543 (1)

VERIFICATION SUMMARY

DATE: 01/27/2003

PATENT APPLICATION: US/10/005,480A

TIME: 10:07:04

Input Set : D:\51158-20062.txt

Output Set: N:\CRF4\01242003\J005480A.raw

L:7450 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:742 after pos.:0
L:8410 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:763 after pos.:0
L:8540 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:765
M:332 Repeated in SeqNo=765